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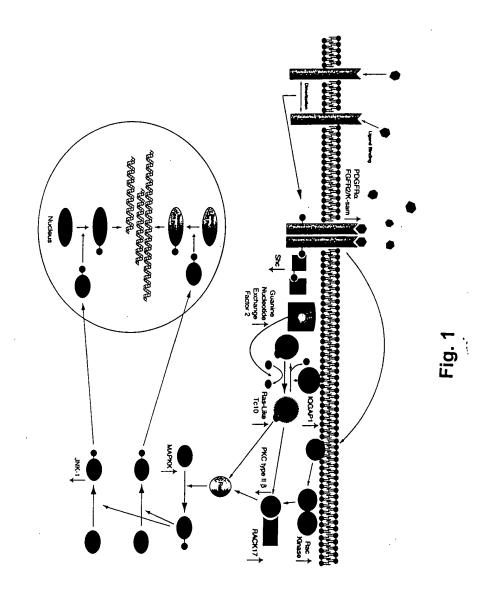
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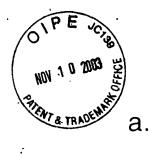
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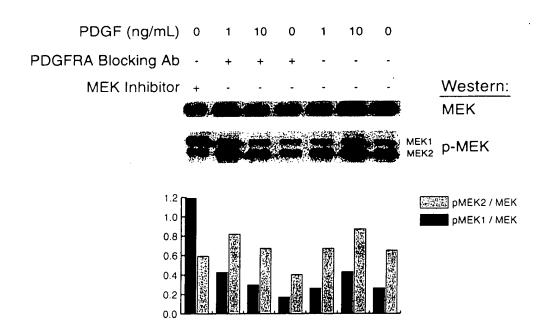
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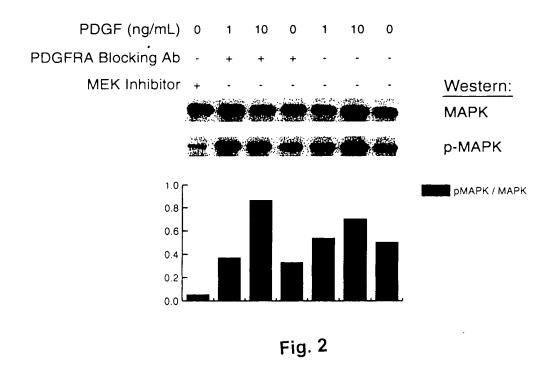








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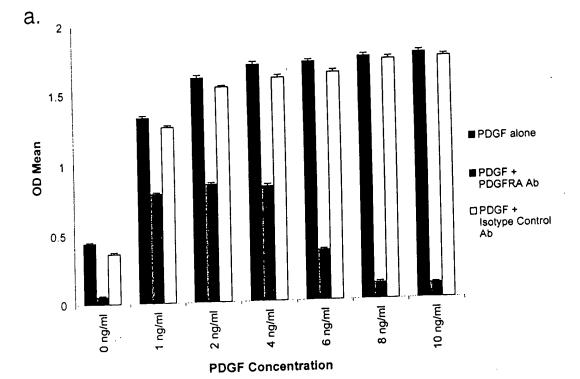


Fig. 3



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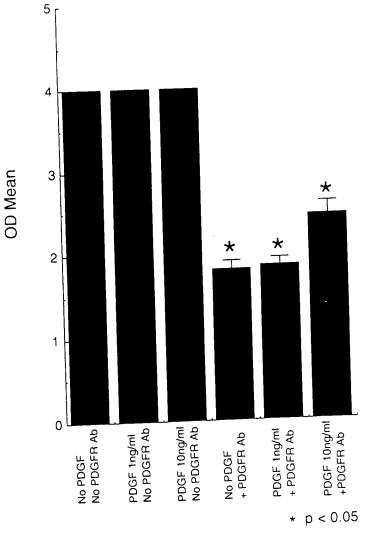
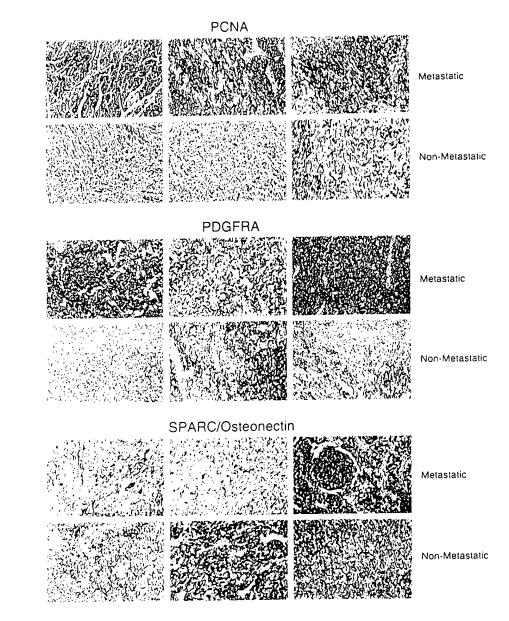


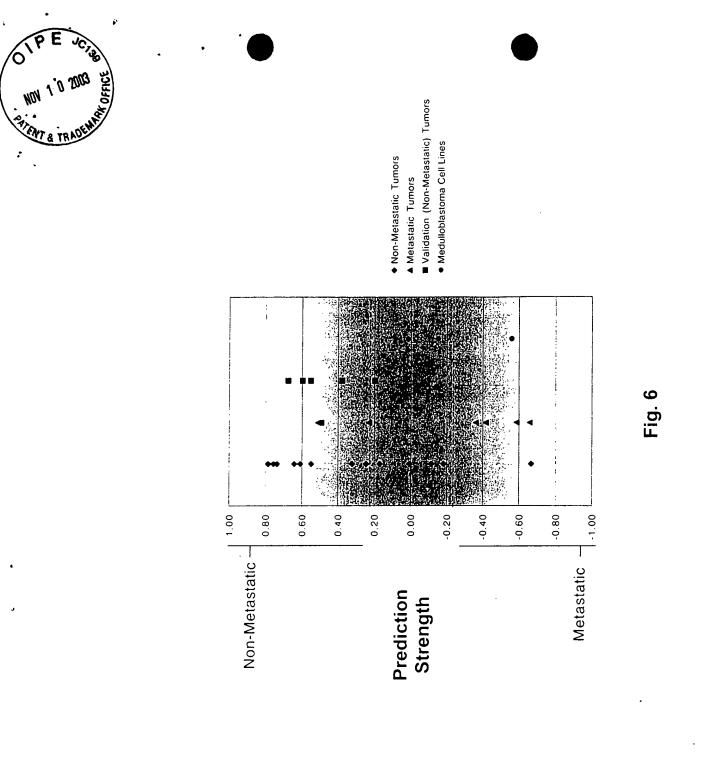
Fig. 4

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Non-Metastatic Tumore	Melasiatic Tumore			Average Intensity, Non-	Average Intensity, Metastatic	Permutational p-value	Avenge Fold Difference	
	,		Gene Name	3404	1896	0.004	1.60	
		424 al	GTP-binding protein (RAB38)	124	23 34	0.006	5.85 2.44	
		1611_6_6 1848_6_6	Interteron (IPN-gamma) Interteution 10 (IL 10)	341	171	0.007	2.25	
		2042_J_N	04740	117	34 297	0.010	1.71	
		199 41	Integrin alpha-3 chain. Human dual-specificity protein phosphatase	1308	440	0.018	2,06	
		1070 Lai	Protein Eness (JHK1) Nedd-4-Bis ubigulin-protein Egass WWP2	241 273	107	0.025	2.64	
		1812_0_4	APC	1567	518 10	0.034	2.65	
		304_81 463_9_8i	Guarina Nucleotide Exchange Factor 2 Nuclear Inctor I B3	3367	1864	0.036	2.04 1.71	
		1380_4	Kansinopyta growth lactor	239 322	140	0.036 0.037	1,76	
·		1800_ai افراقة	Tyronine kinese (TXX) MXII	3264	1812	0.037	1.80	
. a. • ‡		1467_4	Epidermal growth lactor receptor kinase substrate (Epe6) Ribosomal protein S6 kinase 2 (RPS6KAZ)	940 447	471 368	0.040	2.33	
		1127_si 2046_si	Em protein (els-related gene), 3' fank	718	364 80	0.044	1.87	
		2022_4	Rad protein kinase beta Heat shock protein 27 (HSP27)	125	269	0.048	1.82	
14		\$28_41 \$47_8_41	TINUR- HOFI-Brur77 bels-type transcription factor homolog	314	11	0.047	28.22	
		1216_6	Protein tinase C (PKC) type bela 8 p300/CBP-associated factor (P/CAP)	162	12 H	0.048	1.92	_
		1012_44 1611_44	of 2 and p84 isolorms of N-Sht	904	470 437	0.048	1,48	Down
· · · · · · · · · · · · · · · · · · ·		724	Chorlonis Somelomenmotropin Hormone Ce-5 Guarylate kinase associated protein (GKAP)	767 62	26	0.050	2.06	In M+
	L	138_81 808_88	Homeobox 1.4	16	213 11496	6.500	14.40	Upin
	1.1 ·	الر قر 120 افر 120	Oktablione S-transferske-P16 Cathapain D (catD)	2448	6095	6.001	2.11	M+
	<b>2</b> 1	الدورية الم	Rendication protein & 14kDe auburki (RPA)	790 184	1530 3185	0.003	1,84 20,14	
	14 14	1683_8_81 8062_81	Tasue inhibitor of metalloproteinases (HUMTIMP) MAC25	3364	11374	0.004	3.30	
	1.1 1.1	191_41	Mucin (MUC8)	182 217	370 115	0.004	1.92 2.84	
		اه_ا 48 اه_ا 47	Replication protein A 14kDa subunit (RPA) SRARCharaonectin	4166	4584	6.007	2.06	
	· · · · · · · · · · · · · · · · · · ·	1818_at	Res-Like Protein To 10 Insulin-like growth factor binding protein-2	678 417	1318 3012	0.007	4.83	
		اللارية_1741 الدر 1841	Protein kinese C-binding protein RACK17	72	447	0.009	6.19 6.57	
		1321_4_84	Tumor-associated membrane protein homolog (TMP) PGF Receptor K-Barn, All. Splice 3	33 90	343	0.000	3.80	
	i di	لغ_ة_1143 اه_0_1173	Spermidne/Spermine H1-Apetytireneterese, A2, Epilos 2	2401	5486 6071	0.000 0.010	1.48	
	. 1 F	700_sl	Bela-kutulin gene, cione m40 X74784ade receptor protein twosine kinese	3393	408	0.012	8.10	
11		344_41	ST4 Oncoletal antigen	358 281	760 748	0.012	2.16 2.67	
	11	1001_el 942_el	Putative receptor tyrosine tinese (tie) P1-Conte	566	847	0.013	1.82	
11	11	1062_4_41	NF-LE-beta	884 2848	1413	0.013	5.64 1.61	
	14	283_ai 1064_ai	Ubiquinol cytochrome-c reductase core I Replication factor C, 37-kDe suburit	332	\$43	0.015	1.64	
1.2		770_41	Giutathione perceidase	674 513	1879 1451	0.016	2.63	
	11	317_41 1663_4_44	D65695 Cysteine protease Tumor necrosie lactor receptor	928	1675	0.017	1.81 3.64	
		190_41	Likogen induced nuclear orphan receptor (MINOR)	80 1714	2943	0.018	1.67	
		1007_a_ai 1808_ai	Receptor protein-brosine kinase (HEKE)	120	767	0.018 0.019	6.42 1.75	
11	1 <b>1</b> . C	125_81	Gamma-Interteron-Inductble protein (JP-30) Bioom's syndrome protein (BLM)	767 443	1340 896	0.010	1.\$7	
	الفراغة الم	1644_al 816_ <u>0_</u> al	Homeobox protein (HCX7)	10	877 819	0.019	27.71 3.03	
		1137_al	Laukanis virus receptor 2 (GLVR2) RNA polymerase 8 subunit (heRP610)	270 1774	2346	0.021	1.63	
		1306_0_61	Cytophyome P-460LTBV	345 577	641 1444	0.023	1.76	
	1.1	1470_al 1196_al	ONA polymeruse dalla amali subunit RCC1 azonal7-14	220	637	0.025	2.90	
		214_AI	Homeobox protein (HCX7)	1284	2617	0.025	8.04 1.44	
	÷	1782_8_81 736_8_81	Protein Kinase HK31, Camp-Dependent	- 11	129	0.028	12.03	
	11	826_41	(cione 14VS) metallothionein-IG (MT10) mRNA iragment ior beta-3 microglobulin.	212 6916	644 9673	0.028	1.42	
11	1	لمرم_824 . ام_م_115	Fibranectin, All, Ballos 1	661 123	2038	0.030 6.031	3.70 1.67	
		1226_al 1771_4_A	TNF-siphs converting enzyme Platet-derived growth lactor receptor siphs	347	621	9.032	1.62	
	13	1946_1_0	X73064cds NM23-H1	4190	6670	0.033	1.69	
t i		1826_st 1637_st	Rea (GTPease-activating-like protein (IQGAP1) MAPKAP kinese (3pK)	67	172	0.036	1.54	
	່ <u>ປ</u>	له_ا_ 400	Metallothionein I-8	3146 274	4404 881	0.036	1.40 2.40	
	i.3	1970_s_at 1379_at	M58371 Protein tyroeine kinese	346	762	0.037	2.18	
	11	2068_1_6	integrin bela-6 subunit	630 646	1046	0.039	1.66	
		1939_al 1104_a_al	Phosphoprotein p63 Heat shock protein (hep 70)	1983	4117	0.042	2.07	
1.1		133_81	Cathepsin C	442 63	763 257	0.043	1.72 4.62	
	Li	\$024_s_a 702_t_a	: Lyn 8 Homeotic Protein Hps-6	344	618	0.044	1.51	
		892_41	Tumor entigen (L.6)	603	179	0.044	3.36 1.78	
	1	1721_0_6	Mulator gane (NMSH2)	721	1067	0.045	1.47	
		2044.1.4	Alpha1(E)-calenin	2467	3607	0.046	1.42	

# Fig. 7